

DØ New Phenomena Analysis Approach

Greg Landsberg

February 13, 2003

Background

- ◆ New Phenomena group is the leader in terms of the number and the variety of the analyses
 - 7 Analyses for ICHEP 2002
 - 11 Analyses under the EB review for Winter 2003
 - Pretty much all the objects: jets, e , μ , ME_T , τ (!), and b 's (no top yet!)
- ◆ A lot of experience accumulated with various data formats for the past 1.5 years: from root-tuples to TMBTrees
- ◆ This talk reflects our group experience, our praises and our gripes

What's Required for a Good Analysis?

- ◆ No good analysis can be done in a single iteration: multiple data reprocessing is always required
- ◆ An ability to run the analysis code on the highest data tier in real time is imperative
- ◆ An ability to run standard algorithms and corrections at some stage of the analysis is required
- ◆ An ability to trace luminosity information back to RAW data tier is a must
- ◆ Platform-independence of the algorithms and high-level data formats is desirable

New Phenomena Group Data Model

- ◆ We believe that on the long run the TMBTree format fits our analysis model
- ◆ It is reflected in the fact that all 11 analyses are using TMBTree tier
- ◆ It's essential for a successful analysis to use multi-tier data model in order to be able to go significantly far back without reprocessing RAW tier
- ◆ We achieve this via flexible skimming and small analysis samples
- ◆ Disk space is cheap; your time is not!
 - Don't try to save space by dropping chunks – you might regret later
 - Buy disk and fill it; when it's full buy some more or think how to deal with the data which is there
- ◆ In the case of the NP group, we raised \$9K and bought 2.6 TB designated file server: insanity-clued0 with 2x8x180GB RAID arrays and a Gigabit Ethernet

New Phenomena TMB Streams

- TMB is the basic format for our offline skimming

1em	1emtrk	1em1mu	1mutrk
<ul style="list-style-type: none"> 1 EM <p> $ID = 10, 11$ $pT > 20\text{GeV}$ $Iso < 0.15$ $HM < 40$ </p> <p>4.4%</p>	<ul style="list-style-type: none"> 1 EM <p> $ID = 10, 11$ $pT > 12\text{GeV}$ $Iso < 0.15$ </p> <ul style="list-style-type: none"> 1 Track <p> $pT > 7\text{ GeV}$ </p> <ul style="list-style-type: none"> $d\theta < 0.2$ <p>1.3%</p>	<ul style="list-style-type: none"> 1MuonCandidate <p>Medium</p> <ul style="list-style-type: none"> 1 EM <p> $ID = 10, 11$ $pT > 5\text{ GeV}$ </p> <p>2.0%</p>	<ul style="list-style-type: none"> 1MuonCandidate <p>Medium</p> <ul style="list-style-type: none"> 1 Track <p> $pT > 6\text{ GeV}$ </p> <ul style="list-style-type: none"> $d\theta < 0.2$ <p>7.1%</p>

2em	2mu	1em2jets
<ul style="list-style-type: none"> 2 EM <p> $ID = 10, 11$ $pT > 7\text{GeV}$ </p> <p>1.9%</p>	<ul style="list-style-type: none"> 2 MuonCandidate <p>Medium</p> <p>1.0%</p>	<ul style="list-style-type: none"> 1 EM <p> $ID = 10, 11$ $pT > 15\text{GeV}$ $Iso < 0.15$ $HM < 40$ </p> <ul style="list-style-type: none"> 2jets <p> $jetName = JCCA$ $pT > 15\text{ GeV}$ $emETfraction \leq 0.95$ $chETfraction \leq 0.4$ $hotcellratio \leq 10$ $n90 \geq 1$ </p> <ul style="list-style-type: none"> Trigger EM15_2JT15 <p>2.0%</p>

Analysis Samples

- ◆ Every analysis starts with the TMB streams
- ◆ TMB Streams are converted in the TMBTrees, with the latest version (typically 2-3 days per stream)
- ◆ Typically, a preselection code is run on a particular set of the TMBTrees, which allows for a lot of flexibility (can rerun the entire preselection code in less than a day)
- ◆ Currently handicapped by the root bug that does not allow to save a subset of the TMBTree without destroying the cross-reference structure (TObjectArray bug, allegedly fixed in the 3.03.09 root release, to be confirmed)
- ◆ To work around this problem, a few different approaches were used:
 - Create a list of events on the first pass, and then build a tree of these events only, which can be saved further (Alex)
 - Use old root-tuple format instead of TMBTrees (Serban)
 - Run all the corrections/ID routines on the TMBTree and output a very small analysis-specific root-tuple (Yuri)
 - Run all the corrections/ID routines on the TMBTree and output a roottuple, which contains most of the TMBTrees variables, with objects being identified by standard algorithms and references replaced with numerical cross-references (Greg) – this r-tuple can be used by any analysis

Conclusion

- We are generally satisfied with the TMBTree as the basic analysis platform
- We welcome the ID group efforts to make tools that work directly on the TMBTrees (e.g., EMcandidate, Mucandidate, JES)
- More tools of this kind are needed (e.g. ME_T corrections in the rings, etc.)
- Switching from one data format to another is a *major* time sink – please do not do it again!
- We would like the TMBTree format to be debugged and expanded, but we beg that it stays backward compatible and is used as our final attempt of the analysis data tier